



Supplementary file 1

Table S1. Summary of genotyping resistance analysis results.

	Baseline genotype	ART	Virological failure - reason	Genotype at virological failure	Outcome
ART-naïve					
	IN: R263K, T97A; RT: No DRM; PR: No DRM	DTG/abacavir/lamivudine	Week 24 - treatment interruption	Not performed	ART re-initiation and suppression achieved at week 72
	Not performed	EVG/c/tenofovir disoproxyl fumarate/emtricitabine	Week 24 - treatment interruption	IN: E92Q and E157Q RT M184V	N/A
	No mutations	DTG/abacavir/lamivudine	Week 24 - treatment interruption	Not performed	N/A
	No mutations	DTG/abacavir/lamivudine	Week 96 – no reason identified	N/A*	Virological failure at week 96
	No mutations	DTG/abacavir/lamivudine	Week 24 – no reason identified	PR: A71T RT: No DRM IN: N/A*	N/A
	Not performed	DTG/abacavir/lamivudine	Week 24 - suppression not achieved	PR: A71T RT: M184I	Suppression at week 48
	Not performed	EVG/c/emtricitabine/tenofovir disoproxyl fumarate Week 24 - switch to RAL/emtricitabine/tenofovir disoproxyl fumarate**	Week 72 - poor adherence	Not performed	Re-suppression at week 96
ART-switching					
	Not performed	DTG/abacavir/lamivudine	Week 48 – No reason identified	IN: No DRM RT T69N, K70R,	Re-suppression at week 96

ART-salvage

			M184V, T215FV and K219Q PR: No DRM	
IN: T97A; RT: No DRM; PR: No DRM RT: K65R, T69N, V90I, E138A, M184V, M230L IN: not performed PR: L10I; RT: L74I, K103N, E138A, M184V, P225H, K238T IN: No DRM RT: No DRM; PR: No DRM; IN not performed	EVG/c/emtricitabine/tenofovir disoproxyl fumarate DTG/abacavir/lamivudine DTG/abacavir/lamivudine + Darunavir/cobicistat EVG/c/emtricitabine/tenofovir disoproxyl fumarate	Week 48 – No reason identified Week 48 – No reason identified Week 48 - poor adherence Week 24 - treatment interruption	N/A* Not performed IN: E92Q and E157Q PR L10I RT: L74I, K103N, E138A, M184V, P225H, and K238T N/A*	Week 96: HIV-1 RNA of 1130 copies/mL Week 96: HIV-1 RNA of 210 copies/mL. No ART change Re-suppression at week 96 Re-suppression at week 96

DTG, dolutegravir; DRM, drug-related mutations; EVG/c, elvitegravir/cobicistat; IN, integrase gene; N/A, not available; PR, protease gene; RAL, raltegravir; RT, reverse transcriptase gene; ART, antiretroviral treatment.

*Not available due to technical failure of the genotyping procedure (i.e., amplification).

**To avoid potential drug-drug interactions with the treatment for Non-Hodgkin's lymphoma.

Figure 1. Flowchart of study patients

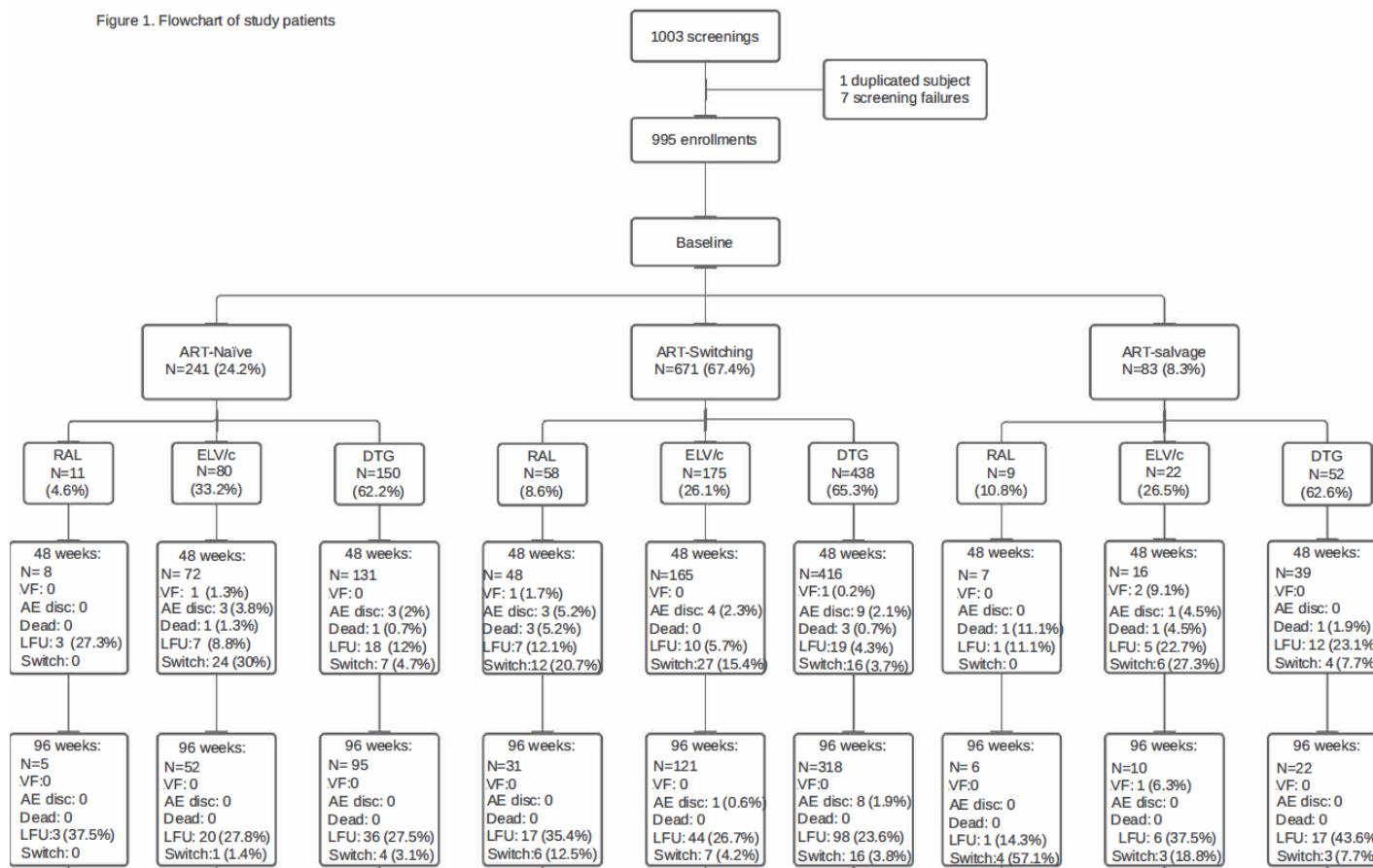
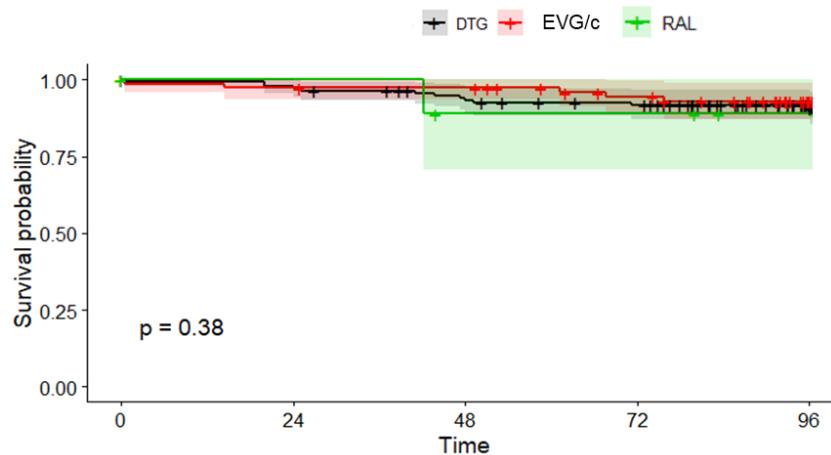


Figure S1. Study flow diagram

A.

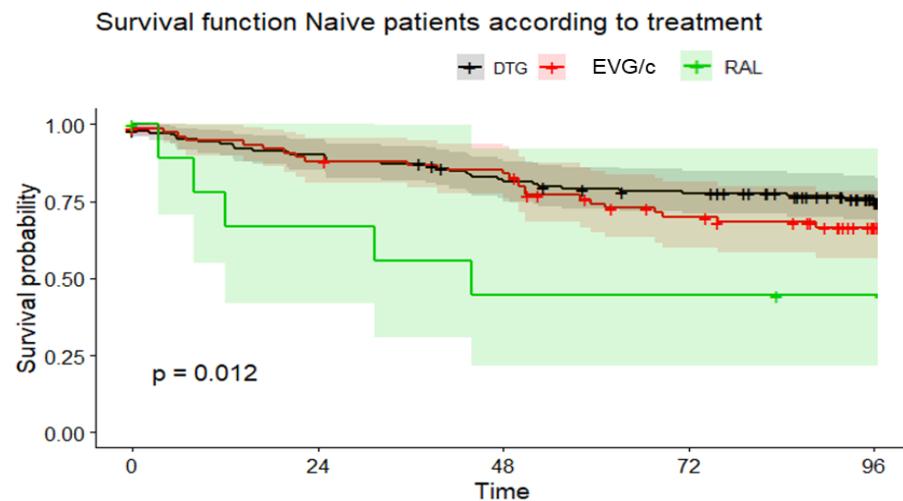
ART-naïve (ITT analysis)



Patients at risk, n	Baseline	24 weeks	48 weeks	72 weeks	96 weeks
DTG	150	135	125	119	79
EVG/c	80	74	68	65	63
RAL	11	11	9	9	3

B.

ART-naïve (ITT sensitivity analysis)

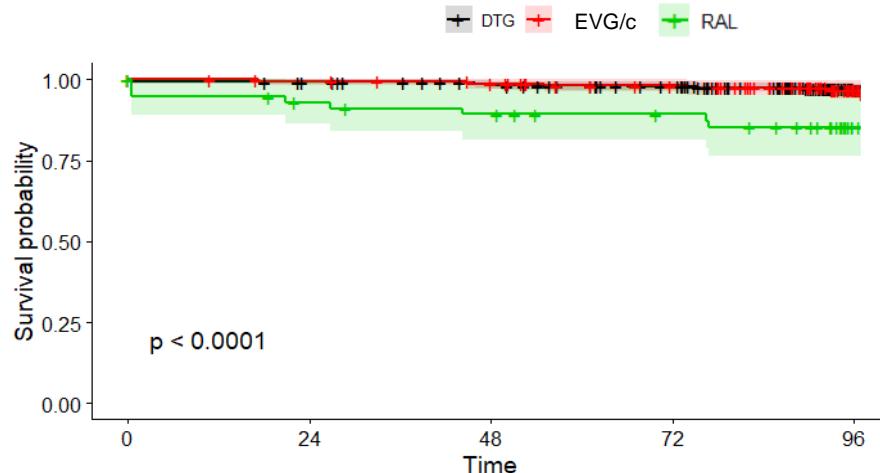


Patients at risk, n	Baseline	24 weeks	48 weeks	72 weeks	96 weeks
DTG	150	127	111	103	69
EVG/	80	67	63	47	39
RAL	11	7	5	5	2

Figure S2. Survival function for virological failure ($CV \geq 50$) in ART-naïve patients according to INSTI-based regimen A. ITT analysis. B. ITT sensitivity analysis. The number of patients at risk at each timepoint are shown below each graph.

A.

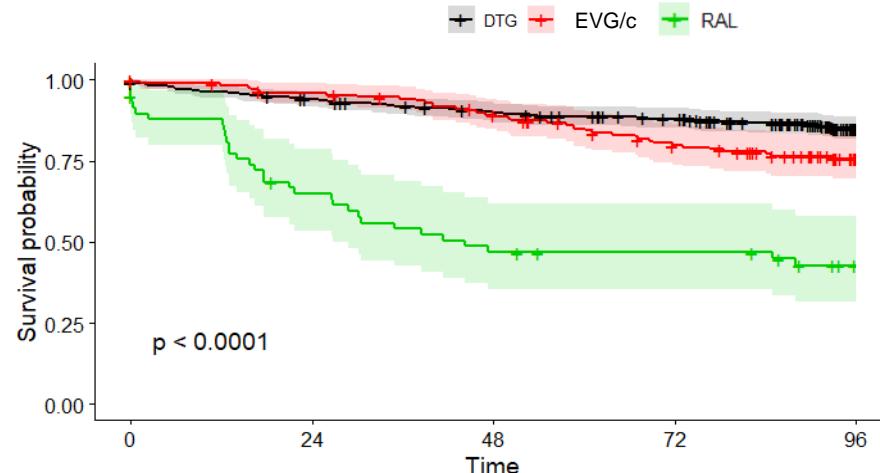
ART-switching (ITT analysis)



Patients at risk, n	Baseline	24 weeks	48 weeks	72 weeks	96 weeks
DTG	438	428	415	403	363
EVG/c	175	168	164	157	129
RAL	58	53	49	44	43

B.

ART-switching (ITT sensitivity analysis)



Patients at risk, n	Baseline	24 weeks	48 weeks	72 weeks	96 weeks
DTG	438	403	381	363	264
EVG/c	175	162	146	126	103
RAL	58	37	27	27	16

Figure S3. Survival function for virological failure (CV \geq 50) in ART-switching patients according to INSTI-based regimen A. ITT analysis. B. ITT sensitivity analysis. The number of patients at risk at each timepoint are shown below each graph.

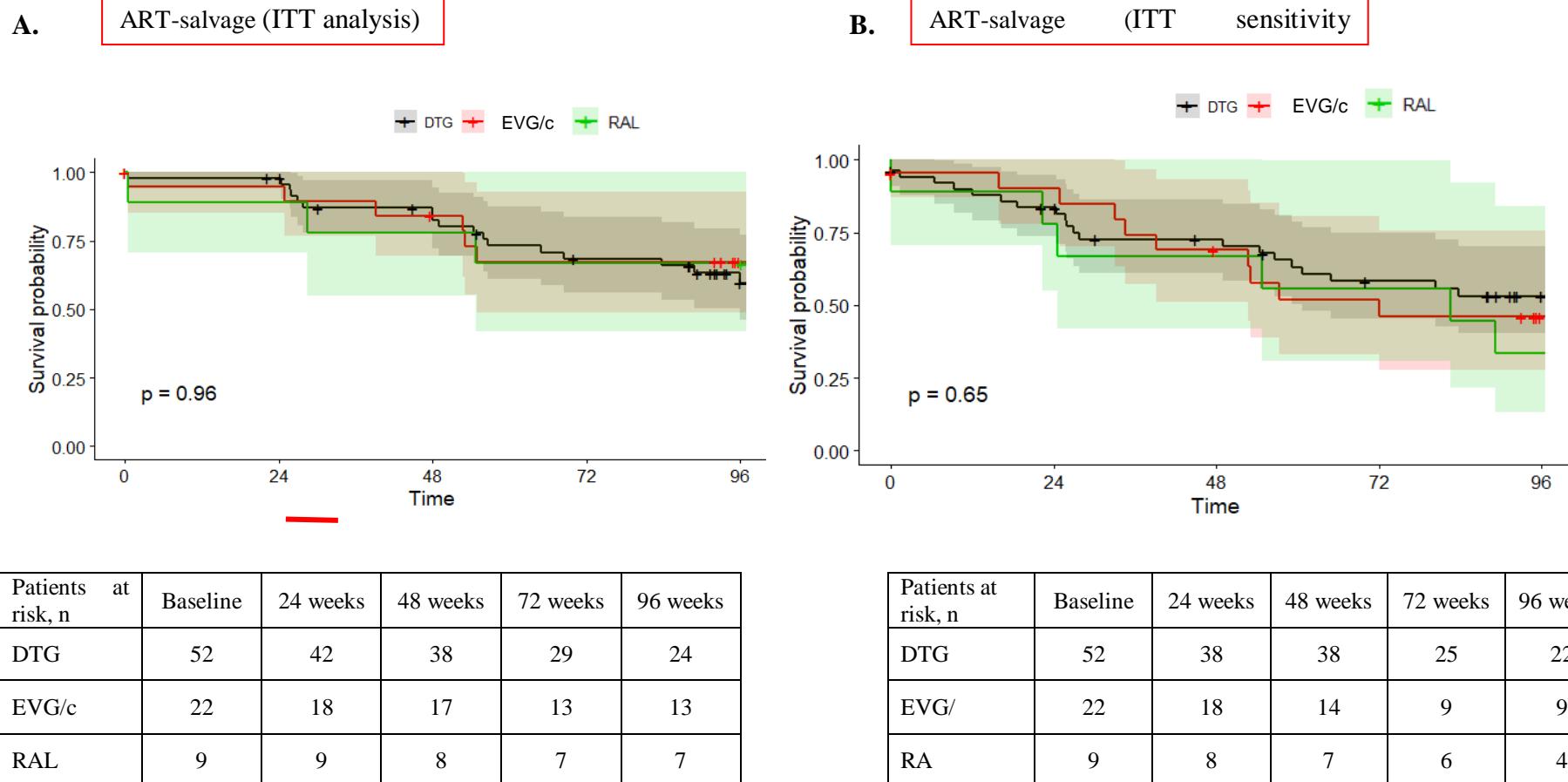


Figure S4. Survival function for virological failure ($CV \geq 50$) in ART-salvage patients according to INSTI-based regimen A. ITT analysis. B. ITT sensitivity analysis. The number of patients at risk at each timepoint are shown below each graph.